

SEQUENCE LISTING

<110> SUGIURA, Takeyuki

<120> TETRAHYDROFOLATE SYNTHETASE GENE

<130> Q94181

<140> US 10/573,969

<141> 2006-03-30

<150> PCT/JP2004/014812

<151> 2004-09-30

<150> JP 2003-341245

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<170> PatentIn version 3.1

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Gln Arg Arg Pro Gln Asp Gly Gln Ala Arg Ser Ser Cys Ser Pro Gly	
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Gly Arg Thr Pro Ala Ala Arg Asp Ser Ile Val Arg Glu Val Ile Gln	
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(1/10)	

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Lys Asp Val Asp Gly Val Thr Asp Ile Asn Leu Gly Lys Leu Val Arg	
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Gln Arg Lys Gly Ser Met Thr Met Ser Ile Gln Trp Lys Thr Arg Gln	
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Leu Gln Ser Lys Leu His Glu Ala Asp Ile Val Val Leu Gly Ser Pro	
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Lys Pro Glu Glu Ile Pro Leu Thr Trp Ile Gln Pro Gly Thr Thr Val	
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cca aga ata cat ttt ggt gga ctc att gag gaa gat gat gtg att ctc	960
Pro Arg Ile His Phe Gly Gly Leu Ile Glu Glu Asp Asp Val Ile Leu	
305 310 315 320	
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325 330 335	
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Trp Leu Arg Glu Gln Gln His Arg Arg Trp Arg Leu His Cys Leu Lys	
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ctt cag cct ctc tcc cct gtg cca agt gac att gag att tca aga gga	1104
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caa act cca aaa gct gtg gat gtc ctt gcc aag gag att gga ttg ctt	1152
Gln Thr Pro Lys Ala Val Asp Val Leu Ala Lys Glu Ile Gly Leu Leu	
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Ser Val Leu Glu Arg Leu Lys Asp Gln Ala Asp Gly Lys Tyr Val Leu	
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Thr Ile Gly Leu Val Gln Ala Leu Thr Ala His Leu Asn Val Asn Ser	
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Phe Ala Cys Leu Arg Gln Pro Ser Gln Gly Pro Thr Phe Gly Val Lys	
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Gly Gly Ala Ala Gly Gly Gly Tyr Ala Gln Val Ile Pro Met Glu Glu	
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Phe Asn Leu His Leu Thr Gly Asp Ile His Ala Ile Thr Ala Ala Asn	
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Val Arg Glu Phe Ser Glu Ile Gln Leu Ala Arg Leu Lys Lys Leu Gly	
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Ile Val Asp Lys Ile Arg Thr Ile Ala Gln Ala Val Tyr Gly Ala Lys	
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Ser Leu Ser His Gln Pro Asp Lys Lys Gly Val Pro Arg Asp Phe Ile	
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Leu Pro Ile Ser Asp Val Arg Ala Ser Ile Gly Ala Gly Phe Ile Tyr	
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Leu Phe	

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45

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Gly Arg Thr Pro Ala Ala Arg Asp Ser Ile Val Arg Glu Val Ile Gln
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Asn Ser Lys Glu Val Leu Ser Leu Leu Gln Glu Lys Asn Pro Ala Phe
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Lys Pro Val Leu Ala Ile Ile Gln Ala Gly Asp Asp Asn Leu Met Gln
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Glu Ile Asn Gln Asn Leu Ala Glu Glu Ala Gly Leu Asn Ile Thr His
 115 120 125

Ile Cys Leu Pro Pro Asp Ser Ser Glu Ala Glu Ile Ile Asp Glu Ile
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Leu Lys Ile Asn Glu Asp Thr Arg Val His Gly Leu Ala Leu Gln Ile
 145 150 155 160

Ser Glu Asn Leu Phe Ser Asn Lys Val Leu Asn Ala Leu Lys Pro Glu
 165 170 175

Lys Asp Val Asp Gly Val Thr Asp Ile Asn Leu Gly Lys Leu Val Arg
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Glu Leu Leu Glu Lys Ser Gly Val Asn Leu Asp Gly Lys Lys Ile Leu
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Val Val Gly Ala His Gly Ser Leu Glu Ala Ala Leu Gln Cys Leu Phe
 225 230 235 240

Gln Arg Lys Gly Ser Met Thr Met Ser Ile Gln Trp Lys Thr Arg Gln
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Lys Pro Glu Glu Ile Pro Leu Thr Trp Ile Gln Pro Gly Thr Thr Val

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 340 345 350

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Gln Thr Asp Lys Ala Leu Tyr Asn Arg Leu Val Pro Leu Val Asn Gly
 (7/10)

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Ala Ser Glu Ile Met Ala Val Leu Ala Leu Thr Asp Ser Leu Ala Asp
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Thr Pro Val Phe Val His Ala Gly Pro Phe Ala Asn Ile Ala His Gly
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Glu Gly Phe Val Val Thr Glu Ala Gly Phe Gly Ala Asp Ile Gly Met
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Gly Pro Ser Val Thr Ala Gly Val Pro Leu Lys Lys Glu Tyr Thr Glu

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755

760

765

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Leu Pro Ile Ser Asp Val Arg Ala Ser Ile Gly Ala Gly Phe Ile Tyr
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Pro Leu Val Gly Thr Met Ser Thr Met Pro Gly Leu Pro Thr Arg Pro
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